(FILE 'HOME' ENTERED AT 12:40:12 ON 18 MAY 2004)

INDEX 'ADISCTI, ADISINSIGHT, ADISNEWS, AGRICOLA, ANABSTR, AQUASCI, BIOBUSINESS, BIOCOMMERCE, BIOSIS, BIOTECHABS, BIOTECHDS, BIOTECHNO, CABA, CANCERLIT, CAPLUS, CEABA-VTB, CEN, CIN, CONFSCI, CROPB, CROPU, DISSABS, DDFB, DDFU, DGENE, DRUGB, DRUGMONOG2, ...' ENTERED AT 12:40:25 ON 18 MAY 2004

SEA HEPARANASE

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FILE 'DGENE, BIOSIS, CAPLUS, SCISEARCH, MEDLINE, EMBASE, USPATFULL, TOXCENTER, CANCERLIT, BIOTECHNO, ESBIOBASE' ENTERED AT 12:41:38 ON 18 MAY

L1

*	2004
L2	174 S L1 AND (SPLICE VARIANT OR HEPARANASE-2)
L3	123 S L2 AND (ISOLAT? OR PURIF?)
L4	123 DUP REM L3 (O DUPLICATES REMOVED)
	FILE 'BIOSIS, CAPLUS, SCISEARCH, MEDLINE, EMBASE, USPATFULL, TOXCENTER,
	CANCERLIT, BIOTECHNO, ESBIOBASE' ENTERED AT 12:47:29 ON 18 MAY 2004
L5	313 S L1 AND (VARIANT OR MUTANT OR SPLICE VARIANT)
L6	225 S L5 AND (PURIF? OR ISOLAT?)
Ь7	1 S L6 AND (HEPARANASE-2)
L8	187 DUP REM L6 (38 DUPLICATES REMOVED)
L9	104 S L1 AND (SPLICE VARIANT)
L10	104 DUP REM L9 (0 DUPLICATES REMOVED)

ANSWER 180 OF 187 CANCERLIT on STN ACCESSION NUMBER: 93696451 CANCERLIT

DOCUMENT NUMBER:

93696451

TITLE:

SOURCE:

The molecular cloning and characterization of human heparanase cDNA and the immunochemical localization

of heparanase in metastatic melanomas.

AUTHOR:

CORPORATE SOURCE:

Univ. of Texas H.S.C. at Houston Grad. Sch. of Biomed. Sci.

Diss Abstr Int [B], (1993) 53 (11) 5515.

ISSN: 0419-4217.

DOCUMENT TYPE:

(THESIS)

LANGUAGE:

English

FILE SEGMENT:

Institute for Cell and Developmental Biology

ENTRY MONTH:

199311

ENTRY DATE:

Entered STN: 19941107

Last Updated on STN: 19970509

Heparanase, an endo-beta-D-glucuronidase, has been associated AB with melanoma metastasis. Polyclonal antibodies directed against the murine N-terminal heparanase peptide detected a Mr of approx 97,000 protein upon SDS-polyacrylamide gel electrophoresis of mouse melanoma and human melanoma cell lysates. In an indirect immunocytochemical study, metastatic human A375-SM and mouse B16-BL6 melanoma cells were stained with the antiheparanase antibodies. Heparanase antigen was localized in the cytoplasm of permeabilized melanoma cells as well as at the cell surface of unpermeabilized cells. Immunohistochemical staining of frozen sections from syngeneic mouse organs containing micrometastases of B16-BL6 melanoma demonstrated heparanase localized in metastatic melanoma cells, but not in adjacent normal tissues. Similar studies using frozen sections of malignant melanomas resected from patients indicated that heparanase is localized in invading melanoma cells, but not in adjacent connective tissues. Monoclonal antibodies directed against murine heparanase were developed and characterized. Monoclonal antibody 10E5, an IgM, precipitated and inhibited the enzymatic activity of heparanase. A 2.6-kb cDNA was isolated from a human melanoma lambda gt11 cDNA library using the monoclonal antibody 10E5. Heparan sulfate cleavage activity was detected in the lysogen lysates from E coli Y1089 infected with the lambda gt11 cDNA and this activity was inhibited in the presence of 10-fold excess of heparin, a potent inhibitor of heparanase. The nucleotide sequence of the cDNA was determined and insignificant homology was found with the gene sequences currently known. The cDNA hybridized to a 3.2-3.4 kb mRNA in human A375 melanoma, WI-38 fibroblast, and THP-1 leukemia cells using Northern blots. Heparanase expression was examined using Western and Northern blots. In comparison to human A375-P melanoma cells, the quantity of 97,000 protein recognized by the polyclonal anti-heparanase antibodies doubled in the metastatic variant A375-SM cells and the quantity of 3.2-3.4 kb mRNA doubled in A375MetMix, a metastatic variant similar to A375-SM cells. In B16 murine melanoma cell, the intensity of the 97,000 protein increased more than 2 times comparing with B16-F1 cells. The extent in the increase of the protein and the mRNA levels is comparable to the change of heparanase activity observed in those cells. In summary, the studies suggest that (a) the N-terminus of the heparanase molecule in mouse and human is antigenically related; (b) heparanase antigens are localized at the cell surface and in the cytoplasm of metastatic human and mouse melanoma cells; (c) heparanase antigens are localized in invasive and metastatic murine and human melanomas in vivo, but not in adjacent normal tissues; (d) heparanase molecule appeared to be differentially expressed at the transcriptional as well as at the translational level; and (e) the size of human heparanase mRNA is 3.2-3.4 kb. (Full text available from University Microfilms International, Ann Arbor, MI, as Order Number AAD93-07237)

ANSWER 187 OF 187 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN

DUPLICATE 9

ACCESSION NUMBER: 1985:362764 BIOSIS

DOCUMENT NUMBER: PREV198580032756; BA80:32756

TITLE: SEQUENTIAL DEGRADATION OF HEPARAN SULFATE IN THE

SUBENDOTHELIAL EXTRACELLULAR MATRIX BY HIGHLY METASTATIC

LYMPHOMA CELLS.

AUTHOR(S): BAR-NER M [Reprint author]; KRAMER M D; SCHIRRMACHER V;

ISHAI-MICHAELI R; FUKS Z; VLODAVSKY I

CORPORATE SOURCE: DEP RADIATION AND CLIN ONCOL, HADASSAH UNIV HOSP, PO BOX

12000, JERUSALEM 91 120, ISRAEL

SOURCE: International Journal of Cancer, (1985) Vol. 35, No. 4, pp.

483-492.

CODEN: IJCNAW. ISSN: 0020-7136.

DOCUMENT TYPE: Article FILE SEGMENT: BA LANGUAGE: ENGLISH

constituents.

A highly metastatic variant (ESb) of a methylcholanthrene-AΒ induced T lymphoma elaborates a heparan sulfate (HS) degrading endoglycosidase (heparanase) to a much higher extent than its non-metastatic parental subline (Eb). Whereas a serum-free medium conditioned by either subline contained a trypsin-like serine protease, heparanase activity was detected only in the ESb-conditioned medium (CM). ESb CM was incubated with a naturally produced, sulfate-labeled subendothelial extracellular matrix (ECM) or with a soluble, high-MW labeled proteoglycan first released from the ECM by incubation with Eb CM or with the partially purified ESb protease. Sulfate labeled degradation products were analyzed by gel filtration on Sepharose 6B. The optimal pH for degradation of ECM-bound HS was 6.2 as compared to pH 5.2 for degradation of the soluble proteoglycan. Heparanase-mediated degradation of both ECM-bound and soluble HS was inhibited by heparin. Addition of either trypsin, plasmin or to a lower extent, the purified ESb protease, stimulated between S- and 20-fold the ESb CM-mediated degradation of ECM-bound HS but had no effect on heparanase-mediated degradation of the soluble proteoglycan. This stimulation was inhibited in the presence of heparin or protease inhibitors. These results indicate that both a protease and heparinase are involved in the ESb-mediated degradation of ECM-bound HS and that 1 enzyme produces a more accessible substrate for the next enzyme. This sequential cleavage is characteristic of degradation of a multimolecular structure such as the subendothelial ECM and hence cannot be detected in studies with its isolated

ANSWER 181 OF 187 BIOSIS COPYRIGHT 2004 BIOLOGICAL ABSTRACTS INC. on STN

DUPLICATE 7

ACCESSION NUMBER: 1994:20409 BIOSIS DOCUMENT NUMBER: PREV199497033409

TITLE:

Nerve growth factor effects on human and mouse melanoma

cell invasion and heparanase production.

Marchetti, Dario [Reprint author]; Menter, Dave; Jin, Li; AUTHOR(S):

Nakajima, Motowo; Nicolson, Garth L.

Dep. Tumor Biol., Box 108, Univ. Texas M.D. Andersen Cancer CORPORATE SOURCE:

Cent., 1515 Holcombe Blvd., Box 108, Houston, TX 77030, USA International Journal of Cancer, (1993) Vol. 55, No. 4, pp.

692-699.

CODEN: IJCNAW. ISSN: 0020-7136.

DOCUMENT TYPE: LANGUAGE:

Article English

ENTRY DATE:

SOURCE:

Entered STN: 25 Jan 1994

Last Updated on STN: 26 Jan 1994

The role of growth factor networks in regulating the progression of human AB melanocytes towards tumorigenicity and ultimately the malignant phenotype is poorly understood. In particular, the autocrine and paracrine influences that modulate cellular invasion and extracellular matrix degradative enzymes of melanoma cells remain undefined at the molecular level. We report here that nerve growth factor (NGF) can modify some metastasis-associated cellular properties of human and mouse melanoma Treatment of early-passage human metastatic melanoma cells (MeWo) or their variants (3S5, 70W) with biologically active 2.5S NGF resulted in (a) delayed density-dependent inhibition of melanoma cell growth; (b) increased in vitro invasion through a reconstituted basement membrane; and (c) time- and dose-dependent induction of heparanase , a heparan-sulfate-specific endo-beta-D-glucuronidase associated with human melanoma metastasis. These effects of NGF were most marked in the 70W brain-colonizing cells (70W gt MeWo gt 3S5). The NGF enhancement of heparanase secretion was not species-specific, since it was also observed in murine B16 melanoma cells; the highest NGF stimulation of heparanase was found in brain-colonizing murine B16-B15b variant (B 16-B15Sb gt B16BL6, B16-F10, B16-F1). NGF also increased the invasive capacity of the human 70W and murine B16-B15b sublines in a chemoinvasion assay performed with filters coated with purified heparan sulfate proteoglycan (HSPG). The enhancement of chemotactic response and heparanase production was detected at NGF concentrations sufficient to fully saturate both low- and high-affinity NGF receptors (NGFR), the neurotrophin receptor (p75) and the trkA gene product, respectively. The results suggest that, in addition to the effects of NGF on cellular development and differentiation within the peripheral and central nervous systems, NGF can exert changes in the invasive properties of neuroectoderm-derived melanoma cells.

L10 ANSWER 104 OF 104 CAPLUS COPYRIGHT 2004 ACS on STN

ACCESSION NUMBER:

2001:472900 CAPLUS

DOCUMENT NUMBER: TITLE:

135:73335 A human heparanase sequence homolog and

splice variants and their possible

therapeutic use in the control of invasive cell

proliferation

INVENTOR(S):

Mckenzie, Edward Alexander; Stamps, Alasdair Craig; Terrett, Jonathan Alexander; Tyson, Kerry Louise

PATENT ASSIGNEE(S):

SOURCE:

Oxford Glycosciences (Uk) Ltd., UK

PCT Int. Appl., 97 pp.

CODEN: PIXXD2

DOCUMENT TYPE:

Patent English LANGUAGE:

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

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PATENT NO.
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        WO 2001046392
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                                     A3
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        WO 2001046392
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              RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, TR, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
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        EP 1240313
                     AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT,
                     IE, SI, LT, LV, FI, RO, MK, CY, AL, TR
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        US 2003083254
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PRIORITY APPLN. INFO.:
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AΒ A human sequence homolog of heparanase and a number of variants that can arise from alternative splicing are described. The protein may play a role in the control of heparan-dependent invasive cell growth in a number of pathologies and may therefore be a target for therapeutics. Identification of an EST for a heparanase homolog in a com. sequence database, PCR cloning of a cDNA and anal. of tissue distribution of the mRNA are reported.

L10 ANSWER 99 OF 104 USPATFULL on STN

ACCESSION NUMBER: 2002:126341 USPATFULL

TITLE: Heparanase II, a novel human

heparanase paralog

INVENTOR(S): Heinrikson, Robert Leroy, Plainwell, MI, UNITED STATES

Bienkowski, Michael Jerome, Portage, MI, UNITED STATES

(9)

NUMBER DATE

PRIORITY INFORMATION: US 2000-199072P 20000420 (60)

DOCUMENT TYPE: Utility FILE SEGMENT: APPLICATION

LEGAL REPRESENTATIVE: Edward F. Rehberg, Pharmacia & Upjohn Company, Global

Intellectual Property, 301 Henrietta Street, Kalamazoo,

MI, 19001

NUMBER OF CLAIMS: 46 EXEMPLARY CLAIM: 1

NUMBER OF DRAWINGS: 4 Drawing Page(s)

LINE COUNT: 2288

CAS INDEXING IS AVAILABLE FOR THIS PATENT.

AB The present invention provides a cDNA encoding a heretofore unknown

enzyme termed heparanase II; constructs and recombinant host cells incorporating the cDNA; the heparanase II polypeptide

encoded by the gene; antibodies to the polypeptide; and methods of

making and using all of the foregoing.